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================================================================
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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2009; month=11; day=18; hr=14; min=30; sec=11; ms=788;]

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Reviewer Comments:

1.

E320   Wrong Nucleic Acid Designator, gc in SEQID (481)

<210> 481

<211> 668

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(668)

* * * * * * * *

His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro
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gtc aca aag agc ttc aac aaa gga gag tgt aag ggc gaa ttc gc   668
Val Thr Lys Ser Phe Asn Lys Gly Glu Cys Lys Gly Glu Phe Ala
    210   215   220

<210> 482

<211> 223

<212> PRT

Codons must be made up of three nucleotides. The last codon in SEQ ID # 481 is only two nucleotides. Please make all necessary changes.

2.

W213   Artificial or Unknown found in <213> in SEQ ID (1)

W213   Artificial or Unknown found in <213> in SEQ ID (2)

W213   Artificial or Unknown found in <213> in SEQ ID (3)

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W213              Artificial or Unknown found in <213> in SEQ ID (4)
W213              Artificial or Unknown found in <213> in SEQ ID (5)
W213              Artificial or Unknown found in <213> in SEQ ID (6)
W213              Artificial or Unknown found in <213> in SEQ ID (7)
W213              Artificial or Unknown found in <213> in SEQ ID (8)
W213              Artificial or Unknown found in <213> in SEQ ID (9)
W213              Artificial or Unknown found in <213> in SEQ ID (10)
W213              Artificial or Unknown found in <213> in SEQ ID (11)
W213              Artificial or Unknown found in <213> in SEQ ID (12)
W213              Artificial or Unknown found in <213> in SEQ ID (13)
W213              Artificial or Unknown found in <213> in SEQ ID (14)
W213              Artificial or Unknown found in <213> in SEQ ID (15)
W213              Artificial or Unknown found in <213> in SEQ ID (16)
W213              Artificial or Unknown found in <213> in SEQ ID (17)
W213              Artificial or Unknown found in <213> in SEQ ID (18)
W213              Artificial or Unknown found in <213> in SEQ ID (19)
W213              Artificial or Unknown found in <213> in SEQ ID (20) This
error has occured more than 20 times, will not be displayed
E257              Invalid sequence data feature in <221> in SEQ ID (636)
E257              Invalid sequence data feature in <221> in SEQ ID (636)
E257              Invalid sequence data feature in <221> in SEQ ID (636)
E257              Invalid sequence data feature in <221> in SEQ ID (637)
E257              Invalid sequence data feature in <221> in SEQ ID (637)
E257              Invalid sequence data feature in <221> in SEQ ID (637)
E257              Invalid sequence data feature in <221> in SEQ ID (637)
E257              Invalid sequence data feature in <221> in SEQ ID (637)


The warnings and errors shown above are ok and require no response.
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Application No: 10045674   Version No: 4.0

Input Set:

Output Set:

Started: 2009-11-05 17:45:21.376
Finished: 2009-11-05 17:45:29.384
Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 8 ms
Total Warnings: 459
Total Errors: 9
No. of SeqIDs Defined: 637
Actual SeqID Count: 637

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
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W 213	Artificial or Unknown found in <213> in SEQ ID (7)
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W 213	Artificial or Unknown found in <213> in SEQ ID (11)
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W 213	Artificial or Unknown found in <213> in SEQ ID (13)
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W 213	Artificial or Unknown found in <213> in SEQ ID (16)
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W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

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              Input Set:


             Output Set:



             Started:   2009-11-05 17:45:21.376
            Finished:   2009-11-05 17:45:29.384
             Elapsed:   0 hr(s) 0 min(s) 8 sec(s) 8 ms
      Total Warnings:   459
        Total Errors:   9
 No. of SeqIDs Defined:   637
    Actual SeqID Count:   637
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Error code	Error Description
	This error has occured more than 20 times, will not be displayed
E 320	Wrong Nucleic Acid Designator, gc in SEQID (481)
E 257	Invalid sequence data feature in <221> in SEQ ID (636)
E 257	Invalid sequence data feature in <221> in SEQ ID (636)
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<110> LADNER, ROBERT C.
      COHEN, EDWARD H.
      NASTRI, HORACIO G.
      ROOKEY, KRISTIN L.
      HOET, RENE
      HOOGENBOOM, HENDRICUS R. J. M.

<120> NOVEL METHODS OF CONSTRUCTING LIBRARIES COMPRISING
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      OF PEPTIDES, POLYPEPTIDES OR PROTEINS AND THE NOVEL
      LIBRARIES

<130> D2033-708931

<140> 10045674
<141> 2001-10-25

<150> 06/198,069
<151> 2000-04-17

<150> 09/837,306
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<160> 637

<170> PatentIn Ver. 3.5

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<210> 2
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<210> 3

<211> 18
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<400> 3
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<210> 7
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<210> 9
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<210> 10
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<210> 11
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<210> 18
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<210> 23

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<400> 24
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<210> 25
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<400> 25
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<210> 27
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<400> 27
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<210> 28
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<400> 28
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<400> 29
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<210> 31
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<210> 32
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<210> 33
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<211> 98
<212> DNA
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<400> 35
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<212> DNA
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<212> DNA
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agatctgagg acacggccgt gtattactgt gcaacaga 98

<210> 39
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<212> DNA
<213> Homo sapiens

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<210> 44
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aaaaccgagg acacggccgt gtattactgt gctagaga 98

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